

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/807,148

DATE: 07/23/2001

TIME: 18:05:42

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07232001\I807148.raw

PS

3 <110> APPLICANT: Tosato, Giovanna et al.  
5 <120> TITLE OF INVENTION: Use of Calreticulin and Calretuculin Fragments to  
6 Inhibit Endothelial Cell Growth and Angiogenesis, and  
7 Suppress Tumor Growth  
9 <130> FILE REFERENCE: 4239 53372  
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/807,148  
C--> 12 <141> CURRENT FILING DATE: 2001-06-04  
14 <150> PRIOR APPLICATION NUMBER: US 60/103,438  
15 <151> PRIOR FILING DATE: 1998-10-06  
17 <160> NUMBER OF SEQ ID NOS: 35  
19 <170> SOFTWARE: PatentIn Ver. 2.0  
21 <210> SEQ ID NO: 1  
22 <211> LENGTH: 1251  
23 <212> TYPE: DNA  
24 <213> ORGANISM: Homo sapiens  
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27 <221> NAME/KEY: CDS  
28 <222> LOCATION: (1)..(1251)  
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32 Met Leu Leu Ser Val Pro Leu Leu Leu Gly Leu Leu Gly Leu Ala Val  
33 1 5 10 15  
35 gcc gag cct gcc gtc tac ttc aag gag cag ttt ctg gac gga gac ggg 96  
36 Ala Glu Pro Ala Val Tyr Phe Lys Glu Gln Phe Leu Asp Gly Asp Gly  
37 20 25 30  
39 tgg act tcc cgc tgg atc gaa tcc aaa cac aag tca gat ttt ggc aaa 144  
40 Trp Thr Ser Arg Trp Ile Glu Ser Lys His Lys Ser Asp Phe Gly Lys  
41 35 40 45  
43 ttc gtt ctc agt tcc ggc aag ttc tac ggt gac gag gag aaa gat aaa 192  
44 Phe Val Leu Ser Ser Gly Lys Phe Tyr Gly Asp Glu Glu Lys Asp Lys  
45 50 55 60  
47 ggt ttg cag aca agc cag gat gca cgc ttt tat gct ctg tcg gcc agt 240  
48 Gly Leu Gln Thr Ser Gln Asp Ala Arg Phe Tyr Ala Leu Ser Ala Ser  
49 65 70 75 80  
51 ttc gag cct ttc agc aac aaa ggc cag acg ctg gtg gtg cag ttc acg 288  
52 Phe Glu Pro Phe Ser Asn Lys Gly Gln Thr Leu Val Val Gln Phe Thr  
53 85 90 95  
55 gtg aaa cat gag cag aac atc gac tgt ggg ggc ggc tat gtg aag ctg 336  
56 Val Lys His Glu Gln Asn Ile Asp Cys Gly Gly Gly Tyr Val Lys Leu  
57 100 105 110  
59 ttt cct aat agt ttg gac cag aca gac atg cac gga gac tca gaa tac 384  
60 Phe Pro Asn Ser Leu Asp Gln Thr Asp Met His Gly Asp Ser Glu Tyr  
61 115 120 125  
63 aac atc atg ttt ggt ccc gac atc tgt ggc cct ggc acc aag aag gtt 432  
64 Asn Ile Met Phe Gly Pro Asp Ile Cys Gly Pro Gly Thr Lys Lys Val  
65 130 135 140  
67 cat gtc atc ttc aac tac aag ggc aag aac gtg ctg atc aac aag gac 480

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Input Set : A:\Pto.amc

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68 His Val Ile Phe Asn Tyr Lys Gly Lys Asn Val Leu Ile Asn Lys Asp  
 69 145 150 155 160  
 71 atc cgt tgc aag gat gat gag ttt aca cac ctg tac aca ctg att gtg 528  
 72 Ile Arg Cys Lys Asp Asp Glu Phe Thr His Leu Tyr Thr Leu Ile Val  
 73 165 170 175  
 75 cgg cca gac aac acc tat gag gtg aag att gac aac agc cag gtg gag 576  
 76 Arg Pro Asp Asn Thr Tyr Glu Val Lys Ile Asp Asn Ser Gln Val Glu  
 77 180 185 190  
 79 tcc ggc tcc ttg gaa gac gat tgg gac ttc ctg cca ccc aag aag ata 624  
 80 Ser Gly Ser Leu Glu Asp Asp Trp Asp Phe Leu Pro Pro Lys Lys Ile  
 81 195 200 205  
 83 aag gat cct gat gct tca aaa ccg gaa gac tgg gat gag cgg gcc aag 672  
 84 Lys Asp Pro Asp Ala Ser Lys Pro Glu Asp Trp Asp Glu Arg Ala Lys  
 85 210 215 220  
 87 atc gat gat ccc aca gac tcc aag cct gag gac tgg gac aag ccc gag 720  
 88 Ile Asp Asp Pro Thr Asp Ser Lys Pro Glu Asp Trp Asp Lys Pro Glu  
 89 225 230 235 240  
 91 cat atc cct gac cct gat gct aag aag ccc gag gac tgg gat gaa gag 768  
 92 His Ile Pro Asp Pro Asp Ala Lys Lys Pro Glu Asp Trp Asp Glu Glu  
 93 245 250 255  
 95 atg gac gga gag tgg gaa ccc cca gtg att cag aac cct gag tac aag 816  
 96 Met Asp Gly Glu Trp Glu Pro Pro Val Ile Gln Asn Pro Glu Tyr Lys  
 97 260 265 270  
 99 ggt gag tgg aag ccc cgg cag atc gac aac cca gat tac aag ggc act 864  
 100 Gly Glu Trp Lys Pro Arg Gln Ile Asp Asn Pro Asp Tyr Lys Gly Thr  
 101 275 280 285  
 103 tgg atc cac cca gaa att gac aac ccc gag tat tct ccc gat ccc agt 912  
 104 Trp Ile His Pro Glu Ile Asp Asn Pro Glu Tyr Ser Pro Asp Pro Ser  
 105 290 295 300  
 107 atc tat gcc tat gat aac ttt ggc gtg ctg ggc ctg gac ctc tgg cag 960  
 108 Ile Tyr Ala Tyr Asp Asn Phe Gly Val Leu Gly Leu Asp Leu Trp Gln  
 109 305 310 315 320  
 111 gtc aag tct ggc acc atc ttt gac aac ttc ctc atc acc aac gat gag 1008  
 112 Val Lys Ser Gly Thr Ile Phe Asp Asn Phe Leu Ile Thr Asn Asp Glu  
 113 325 330 335  
 115 gca tac gct gag gag ttt ggc aac gag acg tgg ggc gta aca aag gca 1056  
 116 Ala Tyr Ala Glu Glu Phe Gly Asn Glu Thr Trp Gly Val Thr Lys Ala  
 117 340 345 350  
 119 gca gag aaa caa atg aag gac aaa cag gac gag gag cag agg ctt aag 1104  
 120 Ala Glu Lys Gln Met Lys Asp Lys Gln Asp Glu Glu Gln Arg Leu Lys  
 121 355 360 365  
 123 gag gag gaa gaa gac aag aaa cgc aaa gag gag gag gag gca gag gac 1152  
 124 Glu Glu Glu Glu Asp Lys Lys Arg Lys Glu Glu Glu Glu Ala Glu Asp  
 125 370 375 380  
 127 aag gag gat gat gag gac aaa gat gag gat gag gag gat gag gag gac 1200  
 128 Lys Glu Asp Asp Glu Asp Lys Asp Glu Asp Glu Glu Asp Glu Glu Asp  
 129 385 390 395 400  
 131 aag gag gaa gat gag gag gaa gat gtc ccc ggc cag gcc aag gac gag 1248  
 132 Lys Glu Glu Asp Glu Glu Glu Asp Val Pro Gly Gln Ala Lys Asp Glu

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133          405          410          415          1251
135 ctg
136 Leu
139 <210> SEQ ID NO: 2
140 <211> LENGTH: 417
141 <212> TYPE: PRT
142 <213> ORGANISM: Homo sapiens
144 <400> SEQUENCE: 2
145 Met Leu Leu Ser Val Pro Leu Leu Leu Gly Leu Leu Gly Leu Ala Val
146   1          5          10          15
148 Ala Glu Pro Ala Val Tyr Phe Lys Glu Gln Phe Leu Asp Gly Asp Gly
149          20          25          30
151 Trp Thr Ser Arg Trp Ile Glu Ser Lys His Lys Ser Asp Phe Gly Lys
152          35          40          45
154 Phe Val Leu Ser Ser Gly Lys Phe Tyr Gly Asp Glu Glu Lys Asp Lys
155          50          55          60
157 Gly Leu Gln Thr Ser Gln Asp Ala Arg Phe Tyr Ala Leu Ser Ala Ser
158   65          70          75          80
160 Phe Glu Pro Phe Ser Asn Lys Gly Gln Thr Leu Val Val Gln Phe Thr
161          85          90          95
163 Val Lys His Glu Gln Asn Ile Asp Cys Gly Gly Gly Tyr Val Lys Leu
164          100         105         110
166 Phe Pro Asn Ser Leu Asp Gln Thr Asp Met His Gly Asp Ser Glu Tyr
167          115         120         125
169 Asn Ile Met Phe Gly Pro Asp Ile Cys Gly Pro Gly Thr Lys Lys Val
170          130         135         140
172 His Val Ile Phe Asn Tyr Lys Gly Lys Asn Val Leu Ile Asn Lys Asp
173   145         150         155         160
175 Ile Arg Cys Lys Asp Asp Glu Phe Thr His Leu Tyr Thr Leu Ile Val
176          165         170         175
178 Arg Pro Asp Asn Thr Tyr Glu Val Lys Ile Asp Asn Ser Gln Val Glu
179          180         185         190
181 Ser Gly Ser Leu Glu Asp Asp Trp Asp Phe Leu Pro Pro Lys Lys Ile
182          195         200         205
184 Lys Asp Pro Asp Ala Ser Lys Pro Glu Asp Trp Asp Glu Arg Ala Lys
185          210         215         220
187 Ile Asp Asp Pro Thr Asp Ser Lys Pro Glu Asp Trp Asp Lys Pro Glu
188   225         230         235         240
190 His Ile Pro Asp Pro Asp Ala Lys Lys Pro Glu Asp Trp Asp Glu Glu
191          245         250         255
193 Met Asp Gly Glu Trp Glu Pro Pro Val Ile Gln Asn Pro Glu Tyr Lys
194          260         265         270
196 Gly Glu Trp Lys Pro Arg Gln Ile Asp Asn Pro Asp Tyr Lys Gly Thr
197          275         280         285
199 Trp Ile His Pro Glu Ile Asp Asn Pro Glu Tyr Ser Pro Asp Pro Ser
200          290         295         300
202 Ile Tyr Ala Tyr Asp Asn Phe Gly Val Leu Gly Leu Asp Leu Trp Gln
203   305         310         315         320
205 Val Lys Ser Gly Thr Ile Phe Asp Asn Phe Leu Ile Thr Asn Asp Glu

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Output Set: N:\CRF3\07232001\I807148.raw

206                    325                    330                    335  
 208 Ala Tyr Ala Glu Glu Phe Gly Asn Glu Thr Trp Gly Val Thr Lys Ala  
 209                    340                    345                    350  
 211 Ala Glu Lys Gln Met Lys Asp Lys Gln Asp Glu Glu Gln Arg Leu Lys  
 212                    355                    360                    365  
 214 Glu Glu Glu Glu Asp Lys Lys Arg Lys Glu Glu Glu Glu Ala Glu Asp  
 215                    370                    375                    380  
 217 Lys Glu Asp Asp Glu Asp Lys Asp Glu Asp Glu Glu Asp Glu Glu Asp  
 218 385                    390                    395                    400  
 220 Lys Glu Glu Asp Glu Glu Glu Asp Val Pro Gly Gln Ala Lys Asp Glu  
 221                    405                    410                    415  
 223 Leu  
 227 <210> SEQ ID NO: 3  
 228 <211> LENGTH: 400  
 229 <212> TYPE: PRT  
 230 <213> ORGANISM: Homo sapiens  
 232 <400> SEQUENCE: 3  
 233 Glu Pro Ala Val Tyr Phe Lys Glu Gln Phe Leu Asp Gly Asp Gly Trp  
 234 1                    5                    10                    15  
 236 Thr Ser Arg Trp Ile Glu Ser Lys His Lys Ser Asp Phe Gly Lys Phe  
 237                    20                    25                    30  
 239 Val Leu Ser Ser Gly Lys Phe Tyr Gly Asp Glu Glu Lys Asp Lys Gly  
 240                    35                    40                    45  
 242 Leu Gln Thr Ser Gln Asp Ala Arg Phe Tyr Ala Leu Ser Ala Ser Phe  
 243                    50                    55                    60  
 245 Glu Pro Phe Ser Asn Lys Gly Gln Thr Leu Val Val Gln Phe Thr Val  
 246 65                    70                    75                    80  
 248 Lys His Glu Gln Asn Ile Asp Cys Gly Gly Gly Tyr Val Lys Leu Phe  
 249                    85                    90                    95  
 251 Pro Asn Ser Leu Asp Gln Thr Asp Met His Gly Asp Ser Glu Tyr Asn  
 252                    100                    105                    110  
 254 Ile Met Phe Gly Pro Asp Ile Cys Gly Pro Gly Thr Lys Lys Val His  
 255                    115                    120                    125  
 257 Val Ile Phe Asn Tyr Lys Gly Lys Asn Val Leu Ile Asn Lys Asp Ile  
 258                    130                    135                    140  
 260 Arg Cys Lys Asp Asp Glu Phe Thr His Leu Tyr Thr Leu Ile Val Arg  
 261 145                    150                    155                    160  
 263 Pro Asp Asn Thr Tyr Glu Val Lys Ile Asp Asn Ser Gln Val Glu Ser  
 264                    165                    170                    175  
 266 Gly Ser Leu Glu Asp Asp Trp Asp Phe Leu Pro Pro Lys Lys Ile Lys  
 267                    180                    185                    190  
 269 Asp Pro Asp Ala Ser Lys Pro Glu Asp Trp Asp Glu Arg Ala Lys Ile  
 270                    195                    200                    205  
 272 Asp Asp Pro Thr Asp Ser Lys Pro Glu Asp Trp Asp Lys Pro Glu His  
 273                    210                    215                    220  
 275 Ile Pro Asp Pro Asp Ala Lys Lys Pro Glu Asp Trp Asp Glu Glu Met  
 276 225                    230                    235                    240  
 278 Asp Gly Glu Trp Glu Pro Pro Val Ile Gln Asn Pro Glu Tyr Lys Gly  
 279                    245                    250                    255

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Input Set : A:\Pto.amc

Output Set: N:\CRF3\07232001\I807148.raw

281 Glu Trp Lys Pro Arg Gln Ile Asp Asn Pro Asp Tyr Lys Gly Thr Trp  
 282 260 265 270  
 284 Ile His Pro Glu Ile Asp Asn Pro Glu Tyr Ser Pro Asp Pro Ser Ile  
 285 275 280 285  
 287 Tyr Ala Tyr Asp Asn Phe Gly Val Leu Gly Leu Asp Leu Trp Gln Val  
 288 290 295 300  
 290 Lys Ser Gly Thr Ile Phe Asp Asn Phe Leu Ile Thr Asn Asp Glu Ala  
 291 305 310 315 320  
 293 Tyr Ala Glu Glu Phe Gly Asn Glu Thr Trp Gly Val Thr Lys Ala Ala  
 294 325 330 335  
 296 Glu Lys Gln Met Lys Asp Lys Gln Asp Glu Glu Gln Arg Leu Lys Glu  
 297 340 345 350  
 299 Glu Glu Glu Asp Lys Lys Arg Lys Glu Glu Glu Glu Ala Glu Asp Lys  
 300 355 360 365  
 302 Glu Asp Asp Glu Asp Lys Asp Glu Asp Glu Glu Asp Glu Glu Asp Lys  
 303 370 375 380  
 305 Glu Glu Asp Glu Glu Glu Asp Val Pro Gly Gln Ala Lys Asp Glu Leu  
 306 385 390 395 400  
 312 <210> SEQ ID NO: 4  
 313 <211> LENGTH: 180  
 314 <212> TYPE: PRT  
 315 <213> ORGANISM: Homo sapiens  
 317 <400> SEQUENCE: 4  
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 319 1 5 10 15  
 321 Thr Ser Arg Trp Ile Glu Ser Lys His Lys Ser Asp Phe Gly Lys Phe  
 322 20 25 30  
 324 Val Leu Ser Ser Gly Lys Phe Tyr Gly Asp Glu Glu Lys Asp Lys Gly  
 325 35 40 45  
 327 Leu Gln Thr Ser Gln Asp Ala Arg Phe Tyr Ala Leu Ser Ala Ser Phe  
 328 50 55 60  
 330 Glu Pro Phe Ser Asn Lys Gly Gln Thr Leu Val Val Gln Phe Thr Val  
 331 65 70 75 80  
 333 Lys His Glu Gln Asn Ile Asp Cys Gly Gly Tyr Val Lys Leu Phe  
 334 85 90 95  
 336 Pro Asn Ser Leu Asp Gln Thr Asp Met His Gly Asp Ser Glu Tyr Asn  
 337 100 105 110  
 339 Ile Met Phe Gly Pro Asp Ile Cys Gly Pro Gly Thr Lys Lys Val His  
 340 115 120 125  
 342 Val Ile Phe Asn Tyr Lys Gly Lys Asn Val Leu Ile Asn Lys Asp Ile  
 343 130 135 140  
 345 Arg Cys Lys Asp Asp Glu Phe Thr His Leu Tyr Thr Leu Ile Val Arg  
 346 145 150 155 160  
 348 Pro Asp Asn Thr Tyr Glu Val Lys Ile Asp Asn Ser Gln Val Glu Ser  
 349 165 170 175  
 351 Gly Ser Leu Glu  
 352 180  
 355 <210> SEQ ID NO: 5  
 356 <211> LENGTH: 61

Use of n and/or Xaa has been detected in the Sequence Listing.  
 Review the Sequence Listing to insure a corresponding  
 explanation is presented in the <220> to <223> fields of  
 each sequence using n or Xaa.

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/807,148

DATE: 07/23/2001

TIME: 18:05:43

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07232001\I807148.raw

L:11 M:270 C: Current Application Number differs, Replaced Application Number  
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:620 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10  
L:629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10  
L:639 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:11  
L:644 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:11  
L:653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11

F05010" 81F40860

PCT

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/807,148

DATE: 07/05/2001

TIME: 15:36:34

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF3\07032001\I807148.raw

**Does Not Comply  
Corrected Diskette Needed**

3 <110> APPLICANT: Tosato, Giovanna et al.  
 5 <120> TITLE OF INVENTION: Use of Calreticulin and Calretuculin Fragments to  
 6 Inhibit Endothelial Cell Growth and Angiogenesis, and  
 7 Suppress Tumor Growth  
 9 <130> FILE REFERENCE: 4239 53372  
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/807,148  
 C--> 12 <141> CURRENT FILING DATE: 2001-06-04  
 14 <150> PRIOR APPLICATION NUMBER: US 60/103,438  
 15 <151> PRIOR FILING DATE: 1998-10-06  
 17 <160> NUMBER OF SEQ ID NOS: 35  
 19 <170> SOFTWARE: PatentIn Ver. 2.0

## ERRORED SEQUENCES

979 <210> SEQ ID NO: 35  
 980 <211> LENGTH: 6  
 981 <212> TYPE: PRT  
 982 <213> ORGANISM: Artificial Sequence  
 984 <220> FEATURE:  
 985 <223> OTHER INFORMATION: Description of Artificial Sequence:Portion of  
 986 retinoic acid receptor  
 988 <400> SEQUENCE: 35  
 989 Lys Gly Phe Phe Arg Arg  
 990 1 5  
 E--> 997 (1)

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/807,148

DATE: 07/05/2001

TIME: 15:36:36

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF3\07032001\I807148.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:620 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10  
L:629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10  
L:639 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:11  
L:644 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:11  
L:653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11  
L:997 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:35

F050703 04F20860



Per/09

Serial Number:

09/807,148

CRF Processing Date:

7/23/2001

Edited by:

Verified by:

(STIC staff)

**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☒ Other: inserted 7" at end of inhibit

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95